

STIC-Biotech/ChemLib

79285

From: Chan, Christina
Sent: Monday, November 04, 2002 7:23 AM
To: Sullivan, Daniel; STIC-Biotech/ChemLib
Subject: RE: Rush Sequence Search 10/080980

Please rush. Thanks Chris

Chris Chan
TC 1600 New Hire Training Coordinator and SPE 1644
308-3973
CM-1, 9B19

-----Original Message-----

From: Sullivan, Daniel
Sent: Saturday, November 02, 2002 11:42 AM
To: Chan, Christina
Subject: Rush Sequence Search 10/080980

Hi Chris,

Here is another sequence search for a 2 month amended case due 11/16.

Please search the standard and issued patent databases for a polynucleotide encoding at least 50 contiguous amino acids of SEQ ID NO:2.

Thank you,
Dan

Daniel M. Sullivan
Examiner AU 1636
Room: 12D12
Mail Box: 11E12
Tel: 703-305-4448

Point of Contact:
Mona Smith
Technical Information Specialist
CM1 6A01
Tel: 308-3278

Searcher: M. Smith
Phone: _____
Location: _____
Date Picked Up: 11/4/02
Date Completed: 11/6/02
Searcher Prep/Review: 5
Clerical: _____
Online time: 5

TYPE OF SEARCH:

NA Sequences: _____
AA Sequences: /
Structures: _____
Bibliographic: _____
Litigation: _____
Full text: _____
Patent Family: _____
Other: _____

VENDOR/COST (where applic.)

STN: _____
DIALOG: _____
Questel/Orbit: _____
DRLink: _____
Lexis/Nexis: _____
Sequence Sys.: _____
WWW/Internet: _____
Other (specify): _____



[illegible]

FEATURES		SOURCE	
<p>Yamashita, K., Kiyosawa, H., Kondo, S., Sakai, T., Shimada, A., Aizawa, K., Kobuchi, S., Hara, A., Itoh, M., Kashi, T., Chikama, K., Akaboshi, T., Ishii, Y., and Hayashizaki, Y.</p> <p>Mapping of young mouse cDNAs to mouse cDNA libraries (1994) <i>FEBS Letters</i> 347: 297-301</p> <p>Figure, Genomics 4, 1994, 272-190 (2-4)</p> <p>Please visit our web site (http://genome.sci.tokyo-u.ac.jp) for further details.</p> <p>c-mouse libraries:</p> <p>Library/Gen/Qualifiers</p> <p>1. 406</p> <p>For primers "Mus musculus"</p> <p>2nd host "Xenopus laevis"</p> <p>3rd host "Xenopus laevis"</p> <p>4th host "Xenopus laevis"</p> <p>5th host "Xenopus laevis"</p> <p>6th host "Xenopus laevis"</p> <p>7th host "Xenopus laevis"</p> <p>8th host "Xenopus laevis"</p> <p>9th host "Xenopus laevis"</p> <p>10th host "Xenopus laevis"</p> <p>11th host "Xenopus laevis"</p> <p>12th host "Xenopus laevis"</p> <p>13th host "Xenopus laevis"</p> <p>14th host "Xenopus laevis"</p> <p>15th host "Xenopus laevis"</p> <p>16th 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$$\begin{aligned} \text{Theorem 1.1.} \quad & \text{Let } \mathcal{H} \text{ be a Hilbert space and } \mathcal{A} \text{ a } \mathcal{C}^* \text{-algebra. Let } \mathcal{K} \text{ be the } \mathcal{C}^* \text{-algebra of compact operators on } \mathcal{H}. \\ & \text{Let } \mathcal{B} \text{ be a } \mathcal{C}^* \text{-algebra and } \mathcal{C} \text{ a } \mathcal{C}^* \text{-algebra. Let } \mathcal{D} \text{ be a } \mathcal{C}^* \text{-algebra. Let } \mathcal{E} \text{ be a } \mathcal{C}^* \text{-algebra. Let } \mathcal{F} \text{ be a } \mathcal{C}^* \text{-algebra.} \end{aligned}$$

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The first of these is the fact that the system is not a simple one. It is a complex system, and the behavior of the system is not predictable. The second is that the system is not a simple one. It is a complex system, and the behavior of the system is not predictable. The third is that the system is not a simple one. It is a complex system, and the behavior of the system is not predictable. The fourth is that the system is not a simple one. It is a complex system, and the behavior of the system is not predictable. The fifth is that the system is not a simple one. It is a complex system, and the behavior of the system is not predictable. The sixth is that the system is not a simple one. It is a complex system, and the behavior of the system is not predictable. The seventh is that the system is not a simple one. It is a complex system, and the behavior of the system is not predictable. The eighth is that the system is not a simple one. It is a complex system, and the behavior of the system is not predictable. The ninth is that the system is not a simple one. It is a complex system, and the behavior of the system is not predictable. The tenth is that the system is not a simple one. It is a complex system, and the behavior of the system is not predictable.

Best Local Similarity:	100.00%	Mismatches:	0
Query Match:	2.77%	Indels:	0
DB:	4	Gaps:	0

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1 TITLE OF INVENTION: A Fast Liquid, A Fragment, Dervoid and DNA
2
3 TITLE & INVENTION: Inventing the Same
4
5 NUMBER OF SEQUENCES: 103
6
7 ADDRESS/NAME ADDRESS:
8
9 STREET: P.O. Box 747
10
11 CITY: Falls Church
12
13 STATE: Virginia
14
15 COUNTRY: USA
16
17 ZIP: 22040-0747
18
19 COMPUTER RELEVABLE FORM:
20
21 MEDIUM TYPE: Floppy disk
22
23 MEDIUM TYPE: floppy disk
24
25 OPERATING SYSTEM: pc dos/MS dos
26
27 OPERATING SYSTEM: pc dos/MS dos
28
29 CURRENT APPLICATION DATA:
30
31 APPLICATION NUMBER: 02/064499.214
32
33 FILING DATE: 10 NOV 1994
34
35 CLASSIFICATION: 45
36
37 ADDRESS/NAME INFORMATION:
38
39 NAME: Murphy Jr., Gerald M.
40
41 REGISTRATION NUMBER: 28,977
42
43 REFERENCE/KEY NUMBER: 1110-100
44
45 TELEPHONE: 703.205.8000
46
47 TELEFAX: 703.205.8000
48
49 INFORMATION FOR SEQ. ID NO.: 41:
50
51 SEQUENCE CHARACTERISTICS:
52
53 LENGTH: 924 base pairs
54
55 TYPE: double-stranded
56
57 STRANDEDNESS: single
58
59 POLARITY: 5' to 3'
60
61 MEDIUM TYPE: DNA
62
63 HYPOTHETICAL: No
64
65 ANTI-SENSE: YES
66
67 ORIGIN: 500bp
68
69 ORGANISM: Human
70
71 US 08 439 214 44
72
73 Alignment Scores:
74
75 Prod. No.: 25.7 Length: 924
76
77 Score: 9.00 Match: 9
78
79 Percent Similarity: 100.000 Conserved: 0
80
81 Percent Similarity: 100.000 Mismatch: 0
82
83 Query Match: 2.77% Gaps: 0
84
85 DB: 4 Gaps: 0
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87 US 10 080 980A 2 (1 425) X US 08 439 214 44 (1 924)
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6
7 OPERATING SYSTEM: pc dos/MS dos
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9 CURRENT APPLICATION DATA:
10
11 APPLICATION NUMBER: 02/064499.214
12
13 FILING DATE: 10 NOV 1994
14
15 CLASSIFICATION: 45
16
17 ADDRESS/NAME INFORMATION:
18
19 NAME: Murphy Jr., Gerald M.
20
21 REGISTRATION NUMBER: 28,977
22
23 REFERENCE/KEY NUMBER: 1110-100
24
25 TELEPHONE: 703.205.8000
26
27 TELEFAX: 703.205.8000
28
29 INFORMATION FOR SEQ. ID NO.: 41:
30
31 SEQUENCE CHARACTERISTICS:
32
33 LENGTH: 924 base pairs
34
35 TYPE: double-stranded
36
37 STRANDEDNESS: not relevant
38
39 POLARITY: not relevant
40
41 MEDIUM TYPE: DNA
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43 HYPOTHETICAL: No
44
45 ANTI-SENSE: No
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47 FEATURES:
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49 NAME/KEY: 5701R
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51 LOCATION: 1..59
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53 FEATURE:
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55 NAME/KEY: 8701R
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57 LOCATION: 906..924
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59 FEATURE:
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61 NAME/KEY: 1..924
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63 LOCATION: 1..924
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65 OTHER INFORMATION: /no/40 "PBI DIT"
66
67 US 08 439 214 45
68
69 Alignment Scores:
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71 Prod. No.: 25.7 Length: 924
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73 Score: 9.00 Match: 9
74
75 Percent Similarity: 100.000 Conserved: 0
76
77 Percent Similarity: 100.000 Mismatch: 0
78
79 Query Match: 2.77% Gaps: 0
80
81 DB: 4 Gaps: 0
82
83 US 10 080 980A 2 (1 425) X US 08 439 214 45 (1 924)
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Page 10

Search completed: November 5, 2002, 21:04:34
Job time : 325 secs

Excerpted from: Molecular Cloning: Laboratory Manual, 2nd Edition, Cold Spring Harbor, NY: Cold Spring Harbor Laboratory Press, 1992.

REFERENCE
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1. (Access 1 to 109201)
Haskins, L. J., Jozwi, M. P., Chatterjee, A., Blumenthal, M.,
Kumar, K. H., and Lander, E. S.

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2. (Access 1 to 109201)
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7. (Access 1 to 109201)
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8. (Access 1 to 109201)
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9. (Access 1 to 109201)
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11. (Access 1 to 109201)
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12. (Access 1 to 109201)
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13. (Access 1 to 109201)
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14. (Access 1 to 109201)
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15. (Access 1 to 109201)
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16. (Access 1 to 109201)
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17. (Access 1 to 109201)
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18. (Access 1 to 109201)
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19. (Access 1 to 109201)
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20. (Access 1 to 109201)
Haskins, L. J., Jozwi, M. P., Chatterjee, A., Blumenthal, M.,
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REFERENCE
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[illegible]

AUTHORS: HUTTON, J.
 TITLE: Direct Submission
 JOURNAL: Submitted (10-m-2001) Sanger Centre, Hinxton, Cambridgeshire,
 CH10 1SA, UK. E-mail: requests.genome@sanger.ac.uk
 COMMENT: requests.genome@sanger.ac.uk
 On Sep 8, 2000 this sequence version replaced 41:6582457
 ----- genome center


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[illegible][illegible]

LOCUS AR017738 566 bp MPNA linear PRI 14-MAR-2000